**(54) Title:** MODIFIED NUCLEOSIDES AND NUCLEOTIDES AND USE THEREOF

**(57) Abstract:** The present invention relates to modified nucleotides and nucleosides and reagents to produce these. The modified nucleotides and nucleotides are assembled to larger oligonucleotides and oligonucleosides, which, for example, may be used for diagnostics of polymorphisms and for antisense therapy of various conditions. The oligonucleotides and oligonucleosides described in the invention have very good endonuclease resistance without compromising the RNA cleavage properties of RNase H wherein combinations of modifications with Y, Z, R or B are claimed: X=O or S, NH or NCH₃, CH₂ Or CH(CH₃), Y=O, S, or NH or NCH₃, CH₂ or CH(CH₃); Z=O, S, or NH or NCH₃, CH₂ or CH(CH₃); R=O or S, or NH or NCH₃, CH₂ or CH(CH₃); B=A, C, G, T; 5-F/cl/BrU or -C, 6-thioguanine, 7-deazaguanine; α- or β-D-(or L)ribo, xylo, arabino or lyxo configuration.

# MODIFIED NUCLEOSIDES AND NUCLEOTIDES AND USE THEREOF

## Field of the invention

The present invention is within the field of molecular biology. More closely, it relates to modified nucleotides and nucleosides and the use thereof as building blocks for incorporation into oligonucleotides and oligonucleosides. These may, for example, be used for antisense therapy.

## Background

The recruitment by RNase H, an endogenous enzyme that specifically degrades target RNA in the antisense oligonucleotide (AON)/RNA hybrid duplex is an important pathway for the antisense action beside the translational arrest. RNase H hydrolyses the RNA strand in an RNA/DNA hybrid in a catalytic manner. It produces short oligonucleotides with 5'-phosphate and 3'-hydroxy groups as final products. Bivalent cations as $Mg^{2+}$ and $Mn^{2+}$ are found to be necessary cofactors for enzymatic activity. The enzyme is widely present in various organisms, including retroviruses, as a domain of the reverse transcriptase. The RNase H1 from *Escherichia coli* is the most characterized enzyme in this family.

RNase H promoted cleavage of the viral mRNA via formation of the duplexes with complementary oligo-DNAs (antisense strand) is one of the strategies to treat pathogen infections and other genetic disorders. Recent isolation of the human RNase H1 and RNase H2 highlights the importance of the development of the antisense drugs utilizing this mechanism of action.

It has been suggested that for eliciting the RNase H in AON/RNA hybrid, the AON part should retain the B-type DNA conformation with 2'-*endo* sugar (South-type, S), while the RNA moiety should retain its A-type helix character with 3'-*endo* sugar (North-type, N). To fulfill these requirements various modifications of sugar, base as well as of the phosphate backbone have been attempted and numerous reports are available about these modified AONs and their antisense action. Among these, AONs having one or more conformationally fixed (either in N- or S-form of the sugar pucker) nucleoside residues have been found to be promising candidates because when they are locked in the N-form, they exhibit high affinity to the target RNA. Recently, the locked nucleic acid (LNA), in

which the sugar moiety is fixed in the North conformation, has shown unprecedented affinity towards RNA. LNA and other modifications which have the fixed N-sugar moiety drive the AON helix to the A-type resulting in RNA/RNA type duplex which accounts for their higher binding affinity, but this leads to the loss of RNase H action. The introduction of conformationally constrained $N$-methanocarba-thymidine residue in the N-form increased the thermodynamic stability of AON/RNA duplex, whereas in the S-form, a destabilizing effect was observed. It was later found that multiple introduction of (N)-methanocarba-thymidines, although increased the thermodynamic stability of the AON/RNA duplex, but failed to recruit any RNase H activity. It is now quite clear that all modifications that lead to preferential North-type sugar, including its constrained form, in an RNA-type AON result in the loss of RNase H activity, because they resemble RNA/RNA duplex, except when they appear at the termini or in the middle in the gapmer-AON. It has been so far assumed that probably three or four N-type conformational repeats are necessary to enhance the thermal stability of RNA-type AON/RNA duplex. Nobody however specifically knows how many North-constrained nucleosides are required to alter the conformational tolerance of the RNase H recognition, thereby its substrate specificity, owing to the *local* structural perturbations in an RNA-type AON/RNA hybrid. On the other hand, 2'-O-methoxy, 2'-F or 2'-O-CH$_2$-CH$_2$-OCH$_3$ based (and other analogous) antisense chemistry, used as a gapmer, promote RNA cleavage by RNase H at least three-fold less satisfactorily than the native. These 2'-O-alkoxy substituted nucleotides are incorporated in the antisense strand as a gapmer to promote complementary RNA cleavage by RNase H. These work better than many other compounds that are available in the literature, but they work less satisfactorily than the native in terms of RNA cleavage efficiency. The efficiency of these 2'-O-methoxy, 2'-F or 2'-O-CH$_2$-CH$_2$-OCH$_3$ based gapmers, "without exception cleaved at slower rate than the wild type substrate" (Crooke et al, *Biochemistry*, **36**, p390-398 (1997)); they work (catalytically) at about 3-fold less efficiency as that of the native counterpart.

Arabino nucleic acids (ANA) have been recently tested for their ability to activate RNase H. Both the sequences tested had lower thermodynamic stability in comparison with the natural DNA/RNA hybrid duplex. CD spectra of these duplexes showed close resemblance to the native DNA/RNA duplexes. Although no quantitative data available, the duplexes formed by ANA and complementary RNA were found to be poorer substrates for RNase H

assisted cleavage compared to the native counterpart. However when $Mn^{2+}$ was used instead of $Mg^{2+}$ in the reaction medium, nearly complete degradation of the target RNA was observed. The 2'F-ANA has also been explored for RNase H potency. Their hybrids with RNA showed higher $T_m$ than the native DNA/RNA hybrid duplex ($\Delta T_m = +5°$ C) and also exhibited global helical conformation similar to native DNA/RNA hybrids as revealed by CD spectroscopy. RNase H promoted cleavage of these 2'F-ANA/RNA hybrids were found to be similar to that observed for native DNA/RNA and DNA-thioate/RNA hybrids. No endonuclease resistance properties of these 2'F-ANA are however known.

Recently, cyclohexenyl nucleosides have been incorporated to AONs (CeNA), and found to have stabilizing effect with the target RNA. The CD spectra of CeNA/RNA hybrid showed close resemblance to the native counterpart. Incorporation of one, two, or three cyclohexenyl-A nucleosides in the DNA strand increases duplex stability with +1.1, +1.6, and 5.2° C. The stabilization effect as expected also depends on the site of introduction. But when tested for RNase H activity they were found to be a relatively poorer substrate for the enzyme in comparison with the native.

Boranophosphate oligothymidines (11mer borano-AON where one of the nonbridging oxygens is replaced with borane) were reported to support RNase H hydrolysis of poly(rA) with efficiency higher than non-modified thymidine oligos regardless of their poor affinity towards the target RNA. The borano modification produces minimal changes in the CD spectrum of the thymidine dimer compared to the native counterpart and both diastereomers adopt B-type conformation (the same as unmodified d(TpT) dimer). Unfortunately, there is no CD or any other structural data available on the hybrid duplexes of such borano-AONs with RNA, which makes it impossible to assess the structural background for the recognition of these duplexes as the substrates by the RNase H vis-à-vis natural counterpart.

Chimeric methylphosphonate based antisense oligos with 5-4-5 methylphosphonates-phosphate- methylphosphonates construct, in particular, having a $T_m$ of about 37°C, was at this temperature more than 4-fold effective at eliciting RNase H hydrolysis of mRNA than the natural congener of $T_m$ 51° C.

## Summary of the invention

The substituted antisense oligonucleotides according to the invention, although show a drop of $T_m$ compared to the native counterpart, can recruit RNase H to cleave the complementary RNA at least as efficiently as the native. The engineering of 3'-exonuclease resistance is rather easily achieved by several means but it is rather difficult to engineer endonuclease resistance without sacrificing on the binding properties to the complementary RNA, or the RNA cleavage by RNase H. The present invention, on the other hand, can combine both of these properties (*i.e.* RNase H mediated cleavage of the complementary RNA strand, as well as the endonuclease resistance of the antisense strand). For example triple oxetane modified oligos show at least four times better endonuclease resistance to the antisense oligos without compromising any RNA cleavage property by RNase H, compared to the native counterpart.

The present inventors have found that the minor groove in AON/RNA duplexes should fulfill following requirements: (1) 1,2-constrained nucleoside derivatives when incorporated in to the AON give the corresrponding AON/RNA duplex preferred helical structure such that the minor groove can accommodate the chemistry of the RNase H cleavage (cleavage site should at least have one B-type DNA conformation in the AON strand with the A-type conformation in the complementary RNA, as suggested by our engineering of the single-point RNA cleavage reaction by RNase H). (2) Such AON/RNA heteroduplexes should be also adequately flexible (as seen by the characteristic lower Tm values, compared to the native counterpart) to accommodate the conformational change required upon complexation with RNAse H – $Mg^{2+}$ in the minor groove for the RNA cleavage by RNase H. (3) The modifications in the minor groove or in its proximity, brought about by a specific 1,2-fused systems in to AON/RNA hybrids do not significantly alter the hydration pattern and secures the availability of the 2'-OH of the RNA for interaction with the active site of RNAse H and $Mg^{2+}$.

In a first aspect, the present invention relates to modified nucleosides and nucleotides, enabling five-membered sugars or their derivatives to be conformationally constrained in the North/East region of the pseudorotational cycle, represented by the following formula:
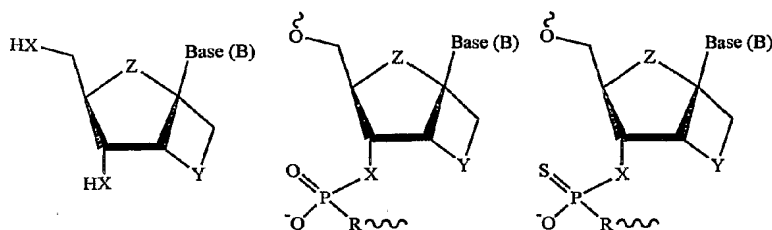
wherein combinations of modifications with X, Y, Z, R or B are claimed:

X = O or S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$),
Y = O, S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$);
Z = O, S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$)
R = O or S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$)
B = A, C, G, T, U, 5-F/Cl/BrU or -C, 6-thioguanine, 7-deazaguanine;

α– or β–D-(or L) ribo, xylo , arabino or lyxo configuration

In a second aspect the invention relates to reagents for the preparation of modified oligonucleotides and oligonucleosides by solid or solution phase synthesis:
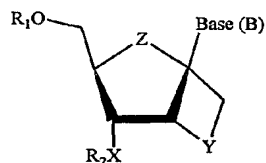


wherein combinations of modifications with Y, Z, R or B are claimed:

X = O or S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$),
Y = O, S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$);
Z = O, S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$)
R = O or S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$)
B = A, C, G, T, U, 5-F/Cl/BrU or -C, 6-thioguanine, 7-deazaguanine;

α– or β–D-(or L) ribo, xylo , arabino or lyxo configuration

R$_1$ = 5'-protecting group according to claim 2.

R$_2$ = 3'-phosphate, 3'-(H-phosphonate), 3'-phosphoramidate, 3'-phosphoramidite, 3'-(alkanephosphonate) according to claim 2.

The different bases, B, may be varied as in claim 2.

In a third aspect, the invention relates to oligonucleotides and oligonucleosides comprising the above modified compounds. These modified monomer blocks according to the invention are introduced (1-9 units) in, for example, antisense oligonucleotides for site-specific modifications, depending upon the length. Thus, the invention provides novels antisense

oligos, AON's. The native nucleotides are fully or partly substituted in the antisense strand by the modified analogs according to the invention.

The oligoribonucleotides and oligoribonucleosides can include substituent groups (both in the tethered and non-tethered form) for modulating binding affinity or artificial nuclease activity to the complementary nucleic acid strand as well as substituent groups for increasing nuclease resistance and for RNase H promoted cleavage of the complementary RNA strand in a site-specific fashion. The oligomeric compounds are useful for assaying for RNA and for RNA products through the employment of antisense interactions, and for the diagnostics, for modulating the expression of a protein in organisms, detection and treatment of other conditions and other research purposes, susceptible to oligonucleotide therapeutics. Synthetic nucleosides and nucleoside fragments are also provided useful for elaboration of oligonucleotides and oligonucleotide analogs for such purposes.

This invention relates for example to compounds based on the oligomeric compounds containing one or more units of 1',2'-fused oxetane, 1',2'-fused azatidine, 1',2'-fused thiatane or 1',2'-fused cyclobutane systems with pentofuranose or the cyclopentane moieties or with any other endocyclic sugar modified (at C4') derivatives ( thereby producing North-East) (N/E) conformationally constrained nucleosides), in either oligonucleotide or oligonucleoside form. These conformationally-constrained nucleosides and nucleotide derivatives (in the N/E constrained structures) in the oligomeric form, when form basepaired hybrid duplexes with the complementary RNA strand, can be useful for modulating the activity of RNA in the antisense therapy or DNA sequencing, in the diagnosis of the postgenomic function or in the design of RNA directed drug development.

In a fourth aspect, the invention relates to therapeutic composition comprising the modified oligonucleotides and oligonucleosides above together with physiologically acceptable carriers.

The main therapeutic use of the composition is antisense therapy of, for example, oncogenic and pathogenic sequences and genetic disorders. Another therapeutic use is to incorporate these blocks into Ribozyme (Catalytic RNA) in order to cleave the target RNA. These blocks can be transformed by nucleoside kinases to the triphosphate form by serving as acceptors

from the phosphate donors such as ATP or UTP (J. Wang, D. Choudhury, J. Chattopadhyaya and S. Eriksson, *Biochemistry*, 38, 16993-16999 (1999). Because of their broader substrate specificities, these triphosphates can interfere with the DNA synthesis of various pathogen and oncogen (antivirals and antitumors).

In a fifth aspect, the invention relates to a diagnostic kit comprising the modified oligonucleotides and oligonucleosides as defined above.

The diagnostic kit is mainly intended for detection of single nucleotide polymorphism SNP and multiple nucleotide polymorphisms MNP. The diagnostic kit is for in vitro use on a human body sample, such as a blood sample. See the following website: http://www.genetrove.com/ of antisense technology for gene functionalization and target validation using 2'-O-alkyl based antisense technology, which is applicable (albeit more efficiently) with the present invention: 1,2-fused sugar technology.

Regulation of how and when genes are turned into proteins can occur at several levels, but RNA is by far the most important generator of complexity and has an enormous potential for creating variation because this go-between molecule stands at the crossroad between genes and proteins. The 1,2-fused system when incorporated in the antisense strand (the antisense technology with the help of RNase H) can be used for systematic studies of how an organism regulates this flexibility through the RNA synthesis and processing (splicing). Thus the antisense technology, using the 1,2-sugar fused nucleoside based chemistries (see the above Figure), is highly relevant to functional genomics - specifically, gene functionalization and target validation, which, in turn to facilitate the discovery and development of new drugs.

In a sixth aspect, the invention relates to a DNA sequencing kit comprising the modified oligonucleotides and oligonucleosides as defined above.

The standard Sanger's dideoxynucleotide sequencing strategy using DNA polymerase and the 2',3'-dideoxynucleotide triphosphates is used (see: http://www.accessexcellence.org/AE/newatg/Contolini/). See also the following website for details of the dideoxynucleotide sequencing strategy: http://www.ultranet.com/~jkimball/BiologyPages/D/DNAsequencing.html

Under the procedure in the website, the 5'-triphosphate building blocks of 1',2'-fused-3'-deoxy- nucleoside (shown below)
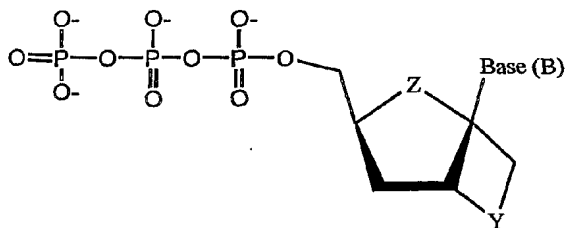
wherein combinations of modifications with Y, Z, or B are claimed:

Y = O, S, or NH or $NCH_3$, $CH_2$ or $CH(CH_3)$;
Z = O, S, or NH or $NCH_3$, $CH_2$ or $CH(CH_3)$
B = A, C, G, T, U, 5-F/Cl/Br-U; 7-deaza-G or hypoxanthine

α– or β–D-(or L) ribo, xylo , arabino or lyxo configuration

are used instead of the standard 2',3'-dideoxynucleotide 5'-triphosphates. The use of 7-deza-guanine or hypoxanthine analog considerably reduce the aggregation owing to the weaker basepairing with dCTP, which, in turn, helps to reduce "compression artifacts" in sequencing gels: http://www.usbweb.com/products/reference/index.asp?Toc_ID=8

In a seventh aspect, the invention relates to use of the modified nucleotides and nucleosides of the invention to produce aptamers (using SELEX procedures, see for example the following website: http://www.somalogic.com/) comprising the modified oligonucleotides and oligonucleosides as defined above. The aptamers may consist of one or several 1,2-modified nucleosides, as defined above, which bind directly to the target proteins or any other ligand, inhibiting their activity.

In an eighth aspect, the invention relates to use of the modified nucleosides, nucleotides and their oligomeric forms of the invention for drug development or in any form of polymerase chain reaction (PCR) or in any molecular biology kit for diagnosis, detection or as reagent.

The present invention was based on the following observations:
1. The introduction of one to five units (North-East) (N/E) conformationally constrained nucleoside(s), such as [1-(1',3'-O-anhydro-β-D-psicofuranosyl)thymine] (T), see claim 1 for

a full list, in to an antisense (AON) strand does not alter the global helical structure of the corresponding AON/RNA hybrid as compared to the native counterpart.

2. Despite the fact that a series of one to five units of N/E-constrained modified AON/RNA hybrid duplexes showed a drop of 2-6°C/modification in $T_m$ (depending upon the number of 1,2-constrained A, C, G or T moieties in the antisense oligo and the composition of sequence), they were cleaved by RNase H with comparable efficiency (or better) as compared to the native counterpart.

3. It was also found that the target RNA strand in the hybrid AON/RNA duplex was resistant up to 5 nucleotides towards 3'-end from the site opposite to the introduction of the N/E-constrained unit in the AON strand, thereby showing the unique transmission of the N/E-constrained geometry of the N/E-constrained residue through the hybrid duplex (*i.e.* the 5-basepaired region has a putative RNA/RNA type duplex structure). An appropriate placement of two such N/E-constrained residues in the AON strand can thus produce a single cleavage site in the complementary RNA strand by RNase H.

4. Despite the fact that some of these sugar-modified AON/RNA duplexes (with three modifications, for example) were destabilized by up to 20 °C compared to the native counterpart, they were found to be as good substrate for RNase H as the native hybrid duplex. The RNase H recruiting power of the oxetane-locked or similarly fused thiatane, azatidine or AONs/RNA hybrids suggests the importance of kinetics as well as relationship between the thermodyanamics of stability/flexibility of hybrid duplexes and the structure/dynamic vis-à-vis recognition, structural tolerance of the hybrid duplex-RNase H complex. Clearly, AON/RNA hybrids should possess certain degree of structural flexibility to undergo certain conformational readjustments upon complexation with RNase H and $Mg^{2+}$ in the minor groove, which is necessary for the cleavage reaction. Those hybrid duplexes which are highly stable have poor conformational flexibility, and are not capable of structurally adjusting themselves upon complexation to the RNase H and $Mg^{2+}$ to form an activated complex to give the cleavage reaction. This is why RNase H do not hydrolyse (or very poorly hydrolyze) those AON/RNA hybrid duplexes which are very stable. Since the RNase H cleavage of the complementary RNA is a slower process than the self-assembly of the AON/RNA hybrid, a smaller population of the hybrid duplex might be actually adequate to bind to RNase H and

drive the complementary RNA cleavage to completion, thereby showing the importance of competing kinetics in the overall cleavage reaction. This is expected to be the case under a non-saturation condition for hybrid duplexes with relatively low $T_m$ as in our oxetane- (or other similarly) modified fused systems.

5. The thermodynamic instabilities of 1,2-fused sugar-modified (*i.e.* N/E-constrained) AONs/RNA hybrids were partially restored by the introduction of dipyridophenazine (DPPZ) moiety at the 3'-end (or at the 5'-end) of these AONs, which also gave enhanced protection towards 3'-exonucleases, and showed equally good RNase H cleavage property as the native counterpart. This was also applied to other 3'-substituents such as cholic acid, folic acid and cholesterol derivatives. All of these tethered substituents were found to be non-toxic in various cellular assays.

6. The loss in the thermodynamic stabilities of 1,2-fused sugar-modified (*i.e.* N/E-constrained) AONs/RNA hybrids with the corresponding oxetane-modified C and G derivatives is ca 2-2.5°C /modification. The actual thermodynamic stability of a given antisense oligo thus depend on the number and type of 1,2-fused sugar-modified A, C, G or T or any other nucleotide blocks

7. The sugar-modified AONs were found to have 3-9 fold more endonuclease resistance compared to those for the native counterparts.

**Detailed description of the invention**

BRIEF DESCRIPTION OF THE DRAWINGS
The numerous objects and advantages of the present invention may be better understood by those skilled in the art by reference to the accompanying figures, in which:

Fig. 1 shows the chemical structure of modified $\underline{T}$ thymine ([1-(1',3'-O-anhydro-β-$\underline{D}$-psico-furanosyl)thymine).

Fig. 2   shows a typical synthetic scheme for the preparation of oxetane-fused nucleosides according to the invention. The following reagents were used: (i) 4-toluoyl chloride, pyridine, r.t., overnight; (ii) silylated base, TMSOTf, acetonitrile, 4° C, 1h, r.t., 18h; (iii) Ms-Cl, pyridine, 4° C, overnight; (iv) 90% aqueous $CF_3COOH$, r.t., 20 min.; (v) NaH, DMF, 4° C, 9h; (vi) methanolic $NH_3$, r.t., 2 days; (vii) DMTr-Cl, pyridine, r.t., overnight;(viii) 2-cyanoethyl-*N,N*-diisopropyl-phosphoramidochloridite, N,N-diisopropylethyl-amine, acetonitrile, r.t., 2h.


The following observations give an insight in to the behavior of various **T** modified AON/RNA hybrids towards RNase H cleavage as well as their stability toward endo and exonucleases:

(1)      The extent of RNA cleavage in hybrid duplexes by *E. coli* RNase H1 in the native hybrid [DNA/RNA] was found to be 68 ± 3%. The target RNA with all single **T**, double **T** and triple **T** modified AONs, were hydrolyzed under the same conditions with extend of 51-68 ± 3%.

(2)      In the AON/ RNA hybrid duplexes with a single mismatch, the RNA was cleaved at a comparable rate as the native counterpart although the hybrid shows a loss of 10 – 11°C in $T_m$ owing to the mismatch. They also showed additional cleavage sites. These two observations therefore show that the recognition of the oxetane-based **T** *vis-a-vis* a mismatch in the AON strand by the target RNA is indeed different, most probably owing to the fact that **T** was perhaps partially hydrogen bonded

(3) The five nucleotide resistance rule to the RNase H cleavage of the RNA in the AON/RNA hybrids in all single **T**, double **T** and triple **T** modified AONs allowed us to engineer a single cleavage site in the target RNA by RNase H. The single RNA cleavage site has been earlier shown to occur in case of 2'-*O*-methyl modified chimeric AON/RNA duplex in which all the central 2'-deoxynucleotides except the middle nucleotide have been shown to adopt an RNA-type conformation by NMR spectroscopy. Since the CD spectra showed that all our **T** modified AON/RNA hybrid duplexes have global structure that corresponds to DNA/RNA type duplex (indicating that our AONs retain the B-DNA type helical conformation in the hybrid), we conclude that the 5-nucleotides resistance rule observed with our **T** modified AONs is owing to more subtle local microscopic conformational (and/or hydration) change, which is only detectable by the enzyme, not by the CD.

(4) The three $\underline{T}$ modified AONs gave the endonuclease stability (with DNase 1) almost 4

fold better (87% of AON remained after 1 h of incubation) compared to the natural

counterpart (19% left), but their 3'-exonuclease stability was identical to that of the native

AON. The 3'-exonuclease stability was however improved by using three $\underline{T}$ modifications

along with the 3'-tethering of dipyridophenazine (DPPZ) moiety, in that 85% of AON was

intact while the native AON was completely hydrolyzed after 2h of incubation with

SVPDE (note that the endonuclease resistance remained however unchanged). The RNase

H promoted cleavage of this AON/RNA duplex (59 ± 4%) remained very comparable to

that of the counterpart with the native AON (68 ± 3%) and with three $\underline{T}$ modified AON (61

± 6%), although a gain of 7°C of $T_m$ was achieved by this additional 3'-DPPZ

modification. This again shows that the rise of $T_m$ do not necessarily dictate the RNase H

cleavage as was earlier found for some methylphosphonate chimeras and

boranophosphates. It should be however noted that the presence of the 3'-DPPZ moiety

produces an additional cleavage site. This is most probably owing to the stabilization of

the terminal G-C hydrogen bonding by the 3'-DPPZ group (observed by NMR) as well as

the recognition of the DPPZ by the enzyme both of which appears to be important for

RNase H recognition, binding and cleavage. Interestingly, amongst all the $\underline{T}$ modified

AONs studied so far, this is the only example where the 5-nucleotide resistance rule in the

RNA strand is not obeyed.


## EXPERIMENTAL PART

**General procedure for preparation of oxetane-modified antisense oligonucleotides (AONs).**

The title compound (**7a**) was prepared from 1,2:3,4-bis-isopropylidene-β-$\underline{D}$ psicofuranose (**1**)

(Fig 2) which was synthesized from $\underline{D}$-fructose. Protection of **1** with 4-toluoyl group to give

**2**, which was coupled with $O,O$-bis(trimethylsilyl)thymine in the presence of TMSOTf as

Lewis acid and acetonitrile as solvent to furnish (1:1) anomeric mixture of the protected

psiconucleosides **3a** (β-isomer) and the corresponding α-isomer in 67% yield. They were

separated by careful column chromatography and the stereochemistry of C2' in **3a** was

confirmed by means of NOE measurements. Methanesulfonylation of β-anomer **3a** afforded

1'-mesylate **4a** (98%) from which the isopropylidine was deprotected using 90% aqueous

$CF_3COOH$ to yield **5a** (92%). The oxetane ring formation was achieved by treatment of **5a**

with NaH in DMF at 0 °C for 9h to give **6a** (60%). Removal of the 4-toluoyl group from **6a**
furnished the desired 1-(1',3'-*O*-anhydro-β-**D**-psicofuranosyl)thymine (**7a**), which was
converted to phosphoramidite building block **9a** (90%) through 6'-*O*-4,4'-dimethoxytrityl
derivative **8a**. The phosphoramidite **9a** was then used for incorporation of **T** residue into
AONs (**3**) – (**6**). Similarly, phosphoramidates **9b** – **10e** were purified and incorporated into
various AONs.

**Typical experiments**

**6'-*O*-4-Toluoyl-1,2:3,4-bis-*O*-isopropyliene-D-psicofuranose (2).** The psicofuranose (**1**)
(5.9 g, 22.5 mmol) was coevaporated with pyridine 3 times and dissolved in 100 ml of the
same solvent. The solution was cooled in an ice bath and 4-toluoyl chloride (3.3 ml, 1.1
mmol) was added dropwise under nitrogen atmosphere. The mixture was stirred at the same
temperature for 2 h. Saturated sodium bicarbonate solution was added and stirring was
continued for further 2 h, and then extracted by DCM. The organic phase was washed with
brine and dried over MgSO$_4$, evaporated and coevaporated with toluene. Recrystallisation
from methanol furnished **2** (7.7 g, 20.2 mmol, 90%). R$_f$ 0.75 (System A). $^1$H-NMR (CDCl$_3$):
7.9 (*d*, J = 8 Hz, 2H), 4-toluoyl; 7.3 (*d*, J = 7.9 Hz, 2H), 4-toluoyl; 4.8 (*d*, J$_{3,4}$ = 5.7 Hz, 1H),
H-4; 4.7 (*d*, 1H), H-3; 4.48-4.35 (*m*, 3H), H-5, H-6, H-6'; 4.33 (*d*, J$_{1,1'}$ = 9.7 Hz, 1H), H-1;
4.1(*d*, 1H), H-1', 2.41 (*s*, 3H), CH$_3$, 4-toluoyl; 1.46 (*s*, 3H), 1.44 (*s*, 3H),1.35,1.33 (*s*, 2×3H)
CH$_3$, isopropyl. $^{13}$C-NMR (CDCl$_3$): 166.3 (C=O, 4-toluoyl); 143.7, 129.8, 128.9, 126.8 (4-
toluoyl); 133.6, 112.7, 111.6; 85.2 (C-3); 82.9 (C-5); 82.3 (C-4), 69.7 (C-1), 64.5 (C-6); 26.4,
26.2, 24.8 (CH$_3$, isopropyl); 21.2 (CH$_3$, 4-toluoyl).

**1-(3',4'-*O*-Isopropyliene-6'-*O*-[4-toluoyl]-α-D-psicofuranosyl)thymine and 1-(3',4'-*O*-
isopropyliene-6'-*O*-[4-toluoyl]-β-D-psicofuranosyl)thymine (3a).** Thymine (3.7 g, 29.6
mmol) was suspended in hexamethyldisilazane (35 ml) and trimethylchlorosilane (5.6 ml)
was added. The reaction mixture was stirred at 120° C in nitrogen atmosphere for 16 h. The
volatile material was evaporated and the residue was kept on an oil pump for 20 min. Sugar **2**
(7.0 g, 18.5 mmol) was dissolved in dry acetonitrile and added to the persilylated nucleobase.
The mixture was cooled to 4° C and trimethylsilyl trifuromethanesulfonate (4.3 ml, 24 mmol)
was added dropwise under nitrogen atmosphere. After being stirred at 4° C for 1h, the
mixture was stirred at room temperature for 18 h. Saturated NH$_4$Cl was added to the reaction
mixture and stirred for 30 min. The organic layer was decanted and the aqueous layer was

extracted 3 times with ether. The combined organic phase was washed first with saturated

sodium bicarbonate solution and then with brine. It was then dried over $MgSO_4$, filtered and

evaporated. The resultant oil was carefully chromatographed using 0-3% MeOH-DCM

yielding **3a** and the corresponding α-anomer. **3a** : (5.5 g, 12.3 mmol, 67%) $R_f$: 0.5 (System

B). (α-anomer of **3a**): $^1$HNMR (CDCl$_3$): 8.8 (*s*, 1H), NH; 7.95 (*d*, J = 8.2 Hz, 2H), 4-toluoyl;

7.5 (*s*, 1H), H-6; 7.28 (*d*, J = 8.4 Hz, 2H), 4-toluoyl; 5.22 (*d*, J$_{3', 4'}$ = 5.9 Hz, 1H), H-3'; 4.83 (*t*,

J$_{4', 5'}$ = 4.7 Hz, 1H), H-4'; 4.71 (*dd*, J$_{gem}$=13.1 Hz, J$_{5', 6'}$ = 7 Hz, 1H), H-6'; 4.55- 4.38 (*m*, 2H),

H-5', H-6''; 4.29 (*dd*, J$_{gem}$= 11.8 Hz, J$_{1', 1'OH}$ = 7.9 Hz, 1H), H-1'; 3.79 (*dd*, J$_{1'', 1'OH}$ = 6.7 Hz,

1H) H-1''; 3.34(*t*, 1H), 1'-OH; 2.43 (*s*, 3H) 4-toluoyl; 1.92 (*s*, 3H), CH$_3$; 1.39, 1.34 (*s*, 2×3H),

CH$_3$. $^{13}$C-NMR (CDCl$_3$): 166.6 (C=O, 4-toluoyl); 164.1 (C-4); 150 (C-2); 144.3 (4-toluoyl);

135.1 (C-6); 129.6, 129.2, 126.2 (4-toluoyl); 113.8 (C-5); 108.9 ( C-Me$_2$); 99.7 (C-2'); 83.7

(C-5'); 82.5 (C-3'); 80.7 (C-4'); 65.1 (C-1'); 63.7 (C-6'); 27, 25.3 (CH$_3$, isopropyl); 21.5

(OCH$_3$); 12.5 (CH$_3$, C-5 CH$_3$). 1D Diff. nOe shows 1.6% nOe enhancement for H6-H5' and

no other nOes expected between other endocyclic-sugar protons and H6 as found for the β-

anomer (see below). (**3a**): $^1$H-NMR (CDCl$_3$): 9.2 (*s*, 1H), NH; 7.71 (*d*, J = 8.2 Hz, 2H), 4-

toluoyl; 7.5 (*s*, 1H), H-6; 7.18 (*d*, J = 7.9 Hz, 2H), 4-toluoyl; 5.44 (*d*, J$_{3', 4'}$ = 6.2 Hz, 1H), H-

3'; 4.87 (*d*, 1H) H-4'; 4.85-4.82 (*m*, 1H), H-5'; 4.65 (*dd*, J$_{gem}$= 12.6 Hz, J$_{5', 6'}$ = 2.4 Hz, 1H), H-

6'; 4.3-4.2 (*m*, J$_{5', 6''}$= 3.7 Hz, 2H), H-6''& H-1'; 3.8 (*dd*, J$_{1'', 1'-OH}$= 6.4 Hz, J$_{gem}$= 12.4 Hz, 1H),

H-1''; 3.27 (*t*, 1H), 1'-OH, 2.4 (*s*, 3H), CH$_3$, 4-toluoyl; 1.6 (*s*, 1H), CH$_3$ (thymine); 1.56, 1.4

(*s*, 2×3H), CH$_3$, isopropyl. $^{13}$C-NMR (CDCl$_3$): 165.6 (C=O, 4-toluoyl); 164.3 (C-4); 150.1 (C-

2); 144.6 (4-toluoyl); 137.3 (C-6); 129.2, 128.9, 125.9 (4-toluoyl); 113.4 (C-5); 108.6 ( C-

Me$_2$), 101.2 (C-2'); 86.1 (C-3'); 83.4 (C-5'); 81.7 (C-4'); 64.2 (C-6'); 63.7 (C-1'); 25.6, 24.1

(CH$_3$, isopropyl); 21.4 (CH$_3$, 4-toluoyl); 11.9 (CH$_3$, thymine). 1D Diff. nOe shows 0.21% nOe

enhancement for H6-H6', 0.08% nOe for H6-H3' and 0.4% nOe for H6-H4' which are

consistent for a β-anomer.

**1-(1'-*O*-Methanesufonyl-3',4'-*O*-isopropyliene-6'-*O*-[4-toluoyl]-β-D-psicofuranosyl)**
**thymine (5a).**Compound **3a** (1.6 g, 3.5 mmol) was coevaporated with pyridine 3 times and

dissolved in 25 ml of the same solvent. The mixture was cooled in an ice bath and

methanesulfonyl chloride (0.75 ml, 9.7 mmol) was added dropwise to the mixture, continued

the stirring for 15 min at the same temperature. The reaction was kept in at 4° C for 12h, then

poured into cold saturated sodium bicarbonate solution and extracted with DCM. The organic

phase was washed with brine, dried over $MgSO_4$, filtered, evaporated and coevaporated with toluene giving compound **5a** (1.89 g, 3.6 mmol, 98%). $R_f$: 0.7 (System B). $^1$H-NMR (CDCl₃): 7.75 (*d*, J = 8.3 Hz, 1H), 4-toluoyl; 7.38 (*d*, J = 1.3 Hz, 1H), H-6; 7.22 (*d*, J = 8.4 Hz, 1H); 4-toluoyl; 5.39 (*d*, $J_{3',4'}$ = 6 Hz, 1H), H-3'; 4.96 (*d*, $J_{gem}$ = 11.4 Hz, 1H), H-1'a; 4.94- 4.88 (*m*, 2H), H-4' & H-5'; 4.7 (*dd*, $J_{gem}$ = 12.6 Hz, $J_{5',6'}$ = 2.5 Hz, 1H); H-6'; 4.39 (*d*, 1H), H-1"; 4.3 (*dd*, $_{5',6''}$ = 3.4 Hz, 1H), H-6"; 2.98 (*s*, 3H), CH₃, OMs; 2.4 (*s*, 3H), CH₃, 4-toluoyl; 1.7, 1.66 (*s*, 2×3H), CH₃, isopropyl. $^{13}$C-NMR (CDCl₃): 165.7 (C=O, 4-toluoyl); 162.9 (C-4); 150.2 (C-2); 145.1 (4-toluoyl); 135.5 (C-6); 129.1, 128.7, 125.6, (4-toluoyl); 114.2 (C-5); 110.1 ( C-Me₂); 98.3 (C-2'); 87.1 (C-3'); 84.2 (C-5' ); 81.7 (C-4'); 69.9 C-1'); 64.1 (C-6'); 37.4 (CH₃, 4-toluoyl); 25.8, 24.3 (CH₃, isopropyl); 21.3 (CH₃, mesyl); 12.3 (CH₃ thymine)


**1-(1'-*O*-Methanesufonyl-6'-*O*-[4-toluoyl]-β-<u>D</u>-psicofuranosyl)thymine (5a)**. Compound **4a** (1.9 g, 3.5 mmol) was stirred with 10.5 ml of 90% CF₃COOH in water for 20 min at r.t. The reaction mixture was evaporated and coevaporated with pyridine. The residue on chromatography furnished **5a** (1.58 g, 3.3 mmol, 92.5%). $R_f$: 0.3 (System B). $^1$H-NMR (CDCl₃ + CD₃OD): 7.75 (*d*, J = 8.3 Hz, 1H), 4-toluoyl; 7.52 (*d*, J = 1.24 Hz, 1H), H-6; 7.2 (*d*, J = 8.4 Hz, 1H), 4-toluoyl; 4.81 (*d*, $J_{gem}$ = 11.6 Hz, 1H), H-1'; 4.76 (*d*, $J_{3',4'}$ = 5.3 Hz, 1H), H-3'; 4.75 (*dd*, $J_{gem}$ = 12.6 Hz, $J_{5',6'}$ = 3.5 Hz, 1H), H-6'; 4.62 (*dt*, 1H), H-5'; 4.58 (*d*, 1H); H-1', 4.41 (*dd*, $J_{4',5'}$ = 3 Hz, 1H), H-4'; 4.33(*dd*, 1H), H-6"; 2.98 (*s*, 3H), CH₃, OMs; 2.4 (*s*, 3H), CH₃, 4-toluoyl; 1.73 (*s*, 3H), CH₃, (thymine). $^{13}$C-NMR (CDCl₃ + CD₃OD): 165.9 (C=O, 4-toluoyl), 163.8 (C-4), 151.7 (C-2); 144.9 (4-toluoyl); 136.3(C-6); 129.2, 129, 126.1 (4-toluoyl); 110.4 (C-5); 97 (C-2'); 83.9 (C-5'); 79.8 (C-3'); 72.2 (C-4'); 69.3 (C-1'); 63 (C-6'); 37.5 (CH₃, 4-toluoyl); 21.3 (CH₃, mesyl); 11.9 (CH₃, thymine)


**1-(1',3'-*O*-Anhydro-6'-*O*-[4-toluoyl]-β-<u>D</u>-psicofuranosyl)thymine (6a)**. To a stirred solution of 80% NaH (171 mg, 5.7 mmol) in 15 ml of DMF in an ice bath, solution of compound **5a** (1.3 g, 2.6 mmol) in 15 ml of DMF was added dropwise. The reaction mixture was stirred at the same temperature for 9h, quenched with 10% acetic acid solution in water and evaporated. The reidue was coevaporated with xylene and on chromatography yielded **6a** (602 mg, 1.5 mmol, 60%). $R_f$: 0.5 (System C). $^1$H-NMR (CDCl₃): 7.93 (*d*, J =8.1 Hz, 2H) 4-toluoyl; 7.25 (*d*, J =7.9 Hz, 2H) 4-toluoyl; 6.81 (*s*, 1H) H-6; 5.47 (*d*, $J_{3',4'}$ = 3.9 Hz, 1H) H-3'; 5.15 (*d*, $J_{gem}$ = 7.9 Hz, 1H) H-1'; 4.79-4.72 (*m*, $J_{gem}$ = 12.3 Hz, $J_{6',5'}$ = 2.55 Hz, 2H) H-1' & H-6'; 4.55-4.42 (*m*, $J_{6'',5'}$ = 2.9 Hz, $J_{4',5'}$ = 8 Hz, 3H), H-4', H-5', H-6"; 2.4 (*s*, 3H), CH₃, 4-

toluoyl, 1.8 (*s*, 3H) CH₃, thymine. ¹³C-NMR (CDCl₃): 166.6 (C=O, 4-toluoyl ), 164.3 (C-4);

149.2 (C-2); 143.8 (4-toluoyl); 135.1 (C-6); 129.5, 128.8, 126.5 (4-toluoyl); 111.6 (C-5); 90.9

(C-2'); 87.3 (C-3'); 80.9 (C-5'); 78.1 (C-1'); 70.3 (C-4'); 63 (C-6'); 21.2 (CH₃, 4-toluoyl); 11.8

(CH₃, thymine)


**1-(1',3'-*O*-Anhydro-β-D̲-psicofuranosyl)thymine (7a).** Compound **6a** (570 mg, 1.5 mmol)

was dissolved in methanolic ammonia (50 ml) and stirred at room temperature for 2 days.

The solvent was evaporated and the residue on chromatography afforded **7a** (378 mg, 1.4

mmol, 96%) $R_f$: 0.3 (System D) ¹H-NMR (CD₃OD, 600 MHz): 7.38 (*d*, J = 1.25 Hz, 1H), H-

6; 5.58 (*d*, $J_{3',4'}$ = 3.8 Hz, 1H), H-3'; 5.33 (*d*, $J_{gem}$ = 8.1 Hz, 1H), H-1'; 4.9 (*d*, 1H), H-1"; 4.46-

4.41(*m*, $J_{4',5'}$ = 8.4 Hz, $J_{5',6'}$ = 2.2 Hz, $J_{5',6"}$ = 5.24 Hz, 2H), H-4'&H-5'; 4.11 (*dd*, $J_{gem}$ = 12.4

Hz, 1H), H-6'; 3.9 (*dd*, 1H), H-6"; 2.1 (*s*, 1H), CH₃, (thymine). ¹³C-NMR (CD₃OD): 166.8 (C-

4); 151.7 (C-2); 138.4 (C-6); 112.7 (C-5); 93.2 (C-2'), 89.3 (C-3'); 85.3 (C-5'); 79.9 (C-1');

71.9 (C-4'); 62.7 (C-6'); 12.1 (CH₃, thymine).


**1-(1',3'-Anhydro-6'-*O*-dimethoxytrityl-β-D̲-psicofuranosyl)thymine (8).** To a solution of

**7a** (353mg, 1.3 mmol) in anhydrous pyridine (6 ml) was added 4,4'-dimethoxytrityl chloride

(510 mg, 1.15 mmol), and the mixture was stirred at r.t overnight. Saturated NaHCO₃

solution was added and extracted with dichloromethane. The organic phase was washed with

brine, dried over MgSO₄, filterd and evaporated. The residue on column chromatography

afforded **8** (647 mg, 1.13mmol, 87%). $R_f$: 0.5 (System B). ¹H-NMR(CDCl3): 7.4-7.1 (*m*,

12H), arom (DMTr)& H-6; 6.85-6.82 (m, 4H), arom (DMTr); 5.4 (*d*, $J_{3',4'}$ = 4.1 Hz, 1H), H-

3'; 5.13 (*d*, $J_{gem}$ = 7.9 Hz, 1H), H-1'; 4.76 (*d*, 1H), H-1"; 4.35 (*dd*, $J_{4',5'}$ = 8.3 Hz, 1H), H-4';

4.28-4.21(*m*, $J_{5',6'}$ = 2.5 Hz, $J_{5',6"}$ = 4.7 Hz, 1H), H-5'; 3.98 (*dd*, $J_{gem}$ = 12.4 Hz, 1H), H-6'; 3.81

(*dd*, 1H), H-6"; 3.8 (*s*, 6H), OCH₃, DMTr; 1.92 (*s*, 3H), CH₃, thymine. ¹³C-NMR (CDCl₃):

164.23, 158.1 (C-4); 149.5; 144.5 (C-2); 135.9, 135.3, 129.8, 128.9, 127.9, 127.5, 126.4,

112.8, (DMTr); 111.6 (C-5); 90.9 (C-2'); 87.6 (C-3'); 83.6 (C-5'); 78.2 (C-1'); 69.7 (C-4');

60.8 (C-6'); 54.9 (DMTr); 11.9 (CH₃, thymine).


**1-(1',3'-Anhydro-6'-*O*-dimethoxytrityl-β-D̲-psicofuranosyl)thymine-4'-*O*-(2-cyanoethyl)-**

**(*N*,*N*-diisopropyl)phosphoramidite (9a).** To a stirred solution of **8** (529 mg, 0.9 mmol) in 5

ml THF, 0.8 ml of *N*,*N*-diisopropyl ethyl amine was added under nitrogen atmosphere and

stirred at r.t for 10 min. To this solution 2-cyanoethyl-*N*,*N*-diisopropyl

phosphoramidochloride (0.4 ml, 1.8 mmol) was added and continued the stirring for 2 h. The reaction was quenched with methanol (3 ml) and the mixture was dissolved in DCM, washed with saturated $NaHCO_3$ solution and brine. The organic layer was dried over $MgSO_4$, filterd and evaporated. The residue on chromatography (30-40% EtOAc, cyclohexane + 2% $Et_3N$) furnished **9a** (632 mg, 0.81 mmol, 90%) $R_f$: 0.5 (system B) The compound was dissolved in DCM 3ml) and precipitated from hexane at -40° C. $^{31}$P-NMR $(CDCl_3)$:150.55; 150.46.


***Synthesis, deprotection and purification of oligonucleotides.*** All oligonucleotides were synthesizesd on 1 μmol scale with 8-channel Applied Biosystems 392 DNA/RNA synthesizer. Synthesis and deprotection of AONs as well as RNA target were performed as previously described.[18] For modified AONs fast depropecting amidites were used and they were deprotected by room temperature treatment of $NH_4OH$ for 16 h. All AONs were purified by reversed-phase HPLC eluting with the following systems: A (0.1 M triethylammonium acetate, 5% MeCN, pH 7) and B (0.1 M triethylammonium acetate , 50% MeCN, pH 7). The RNA target was purified by 20% 7 M urea polyacrylamide gel electrophoresis and its purity and of all AONs (greater than 95%) was confirmed by PAGE. Representive data from MALDI-MS analysis: AON (4) [M-H]⁻ 4478.7; calcd 4478; RNA target (7) [M-H]⁻ 4918.1; calcd 4917.1.


### 1-(1',3'-*O*-Anhydro-β-D-psicofuranosyl)uracil (7b)

$^1$H-NMR($CD_3OD$): 7.48 (d, $J_{5,6}$ = 8 Hz, 1H, H-6), 5.81(d, 1H, H-5), 5.49 (d, $J_{3', 4'}$ = 3.1 Hz, 1H, H-3'), 5.24 (d, $J_{gem}$ = 8 Hz, 1H, H-1'), 4.8 (d, 1-H, H-1"), 4.38-4.3 (m, $J_{4',5'}$ = 8.1 Hz, $J_{5',6'}$ = 1.6 Hz, $J_{5',6"}$ = 6 Hz, 2H, H-4' and H-5'), 4.04 (dd, $J_{gem}$ = 13 Hz, 1H, H-6'), 3.83 (dd, 1H, H-6"). $^{13}$C-NMR $(CD_3OD)$: 166.4 (C-4), 151.4 (C-2), 143 (C-6), 103.6 (C-5), 93 (C-2'), 89.3 (C-3'), 85.4 (C-5'), 79.9 (C-1'), 71.8 (C-4'), 62.6 (C-6').


### 1-(1',3'-*O*-Anhydro-β-D-psicofuranosyl)cytosine (7c).

$^1$H-NMR($D_2O$): 7.28 (d, $J_{5,6}$ = 7.3 Hz, 1H, H-6), 5.94 (d, 1H, H-5), 5.44 (d, $J_{3', 4'}$ = 3.1 Hz, 1H, H-3'), 5.14 (d, $J_{gem}$ = 8.3 Hz, 1H, H-1'), 4.76 (d, 1-H, H-1"), 4.29-4.23 (m, $J_{5',6"}$ = 4.9 Hz, 2H, H-4' and H-5'), 3.9 (d, $J_{gem}$ =12.3 Hz, 1H, H-6'), 3.74 (dd, 1H, H-6"). $^{13}$C-NMR $(D_2O)$: 166.5 (C-4), 156.1 (C-2), 141.9 (C-6), 96.4 (C-5), 91.8 (C-2'), 87.5 (C-3'), 82.6 (C-5'), 78.7 (C-1'), 69.6 (C-4'), 60.5 (C-6').

**RNase H digestion assays**

DNA/RNA hybrids (0.8 μM) consisting of 1:1 mixture of antisense oligonucleotide and target RNA (specific activity 50000 cpm) were digested with 0.3 U of RNase H in 57 mM Tris-HCl (pH 7.5), 57 mM KCl, 1 mM $MgCl_2$ and 2mM DTT at 21-37 °C. Prior to the addition of the enzyme reaction components were preannealed in the reaction buffer by heating at 80 °C for 4 min followed by 1.5 h. equilibration at 21-37 °C. Total reaction volume was 26 μl. Aliquots (7 μl) were taken after 5, 15, 30, 60 and 120 min and reaction was stopped by addition of the equal volume of 20 mM EDTA in 95% formamide. RNA cleavage products were resolved by 20% polyacrylamide denaturing gel electrophoresis and visualized by autoradiography. Quantitation of cleavage products was performed using a Molecular Dynamics PhosphorImager. The experiment is repeated at least 4 times and average values of the % of cleavage are reported here.

**Exonuclease degradation studies**

Stability of the AONs towards 3'-exonucleases was tested using snake venom phosphodiesterase from *Crotalus adamanteus*. All reactions were performed at 3 μM DNA concentration (5'-end ^{32}P labeled with specific activity 50000 cpm) in 56 mM Tris-HCl (pH 7.9) and 4.4 mM $MgCl_2$ at 22 °C. Exonuclease concentration of 70ng/μl was used for digestion of oligonucleotides (total reaction volume was 16μl). Aliquots were quenched by addition of the same volume of 20 mM EDTA in 95% formamide. Reaction progress was monitored by 20% 7 M urea PAGE and autoradiography.

**Endonuclease degradation studies**

Stability of AONs towards endonuclease was tested using DNase 1 from *Bovine pancreas*. Reactions were carried out at 0.9 μM DNA concentration (5'-end ^{32}P labeled with specific activity 50 000 cpm) in 100mM Tris-HCl (pH 7.5) and 10 mM $MgCl_2$ at 37 °C using 30 unit of DNase 1 (total reaction volume was 22μl). Aliquots were taken at 60, 120, 180 and 240 min and quenched with the same volume of 20 mM EDTA in 95% formamide. They were resolved in 20% polyacrylamide denaturing gel electrophoresis and visualized by autoradiography.

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# CLAIMS

1. Modified nucleosides and nucleotides, represented by the following formula:
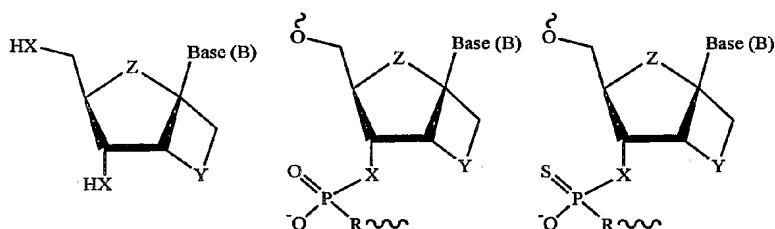


wherein combinations of modifications with X, Y, Z, R or B are claimed:

X = O or S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$),
Y = O, S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$);
Z = O, S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$)
R = O or S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$)
B = A, C, G, T, U, 5-F/Cl/BrU or -C, 6-thioguanine, 7-deazaguanine;

α– or β–D-(or L) ribo, xylo, arabino or lyxo configuration

and oligonucleotides and oligonucleosides comprising these.

2. Reagents for the preparation of modified nucleoside-nucleotide analogs, oligonucleotides or oligonucleosides by solid or solution phase synthesis:

wherein combinations of modifications with Y, Z, R or B are claimed:

X = O or S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$),
Y = O, S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$);
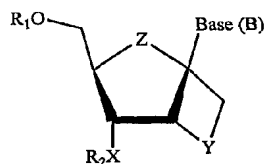Z = O, S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$)
R = O or S, or NH or NCH$_3$, CH$_2$ or CH(CH$_3$)
B = A, C, G, T, U, 5-F/Cl/BrU or -C, 6-thioguanine, 7-deazaguanine;

α– or β–D–(or L) ribo, xylo , arabino or lyxo configuration


comprising the following possible variations of the building blocks for synthesis of compounds:.


for Uracil:

N3-Benzoyl, N3-(4-toluoyl), N3-(2-toluoyl), N3-(4-anisoyl); N3-(4-chlorobenzoyl), N3 - (2,2,2-trichloro-t-butyloxycarbonyl), N3 -(triphenylmethanesulfenyl), N3-(butylthio-carbonyl), N3-(methoxyethoxymethyl), O4-(2-Nitrophenyl), O4-(2-(4-cyanophenyl)-ethyl), O4-(2-(4-nitrophenyl)-ethyl), O4-phenyl, O4-2-methylphenyl, O4-(4-methylphenyl), O4-(2,4-di-methylphenyl), O4-(3-chlorophenyl), O4-(2-(4-nitrophenylsulfonyl)-ethyl), O4-(6-methyl-3-pyridyl), O4-(4-nitrophenylethoxycarbonyl), O4-(4-methyl-3-pyridyl), O4-2,4,6-trimethylphenyl,


for Cytosine:

N4-Anisoyl, N4-benzoyl, N4-(3,4-dimethylbenzoyl), N4-acetyl, N4-phenoxyacetyl, N4-dimethylaminomethylene, N4-benzyloxycarbonyl, N4-levulinoyl, N4-isobutyryl, N4-(2-nitrophenylsulfenyl), N4-isobutoxycarbonyl, N4-(2,2,2-trichloro-t-butyloxycarbonyl), N4-(9-fluorenylmethoxycarbonyl), N4-(N-methyl-2-pyrrolidine amidine), N4-(N,N-di-n-butylformamidine), N4-(3-methoxy-4-phenoxybenzoyl), N4-(isopropoxyacetyl), N4-(2-(tertbutyldiphenylsilyloxymethyl)-benzoyl), N4-(phenylsulfonylethoxycarbonyl), N4-(4-chlorophenylsulfonylethoxycarbonyl), N4-(2-chlorophenylsulfonylethoxycarbonyl), N4-(4-nitrophenylethoxycarbonyl), N4-2-(acetoxymethyl)benzoyl,

for Adenine:

N6-Di-n-butylformamidine, N6-benzoyl, N6-succinyl, N6, N6-phthaloyl, N6-(4,5-
dichlorophthaloyl), N6-tetrachlorophthaloyl, N6-(2-(4-nitrophenyl)-ethoxycarbonyl), N6-
phenoxyacetyl, N6-(9-fluorenylmethoxycarbonyl), N6-(3-chlorobenzoyl), N6-anisoyl), N6-
(4-tertbutylbenzoyl), N6-phenoxycarbonyl, N6-benzyloxycarbonyl, N6-isobutoxycarbonyl,
N6-(2,2,2-trichloro-t-butyloxycarbonyl), N6-dimethylacetamidine, N6-(2-
nitrophenylsulfenyl), N6-dimethylaminomethylene, N6-di-n-butylaminomethylene, N6-(N-
methyl-2-pyrrolidine amidine), N6-(N,N-di-n-butylformamidine), N6-(3-methoxy-4-
phenoxybenzoyl), N6-isopropoxyacetyl, N6-(2-(tertbutyldiphenylsilyloxymethyl)-benzoyl),
N6-phenylsulfonylethoxycarbonyl, N6-(4-chlorophenylsulfonylethoxycarbonyl), N6-(2-
chlorophenylsulfonylethoxycarbonyl), N6-(4-nitrophenylethoxycarbonyl), N6-2-
(acetoxymethyl)benzoyl, N6-(m-chlorobenzoyl).


for Guanine and 7-deazaguanine (hypoxanthine has the same O6 protection as guanine or 7-
deazaguanine):

N2 -Isobutyryl, acetyl, N2-(4-tertbutylbenzoyl), N2-benzyloxycarbonyl, N2-phenoxyacetyl,
N2-benzoyl, N2-levulinoyl, N2-(2-nitrophenylsulfenyl), N2-(9-fluorenylmethoxycarbonyl),
N2-(2,2,2-trichloro-t-butyloxycarbonyl), N2-propionyl, N2-dimethylaminomethylene, N2-
dimethylacetamidine, N2-(N-methyl-2-pyrrolidineamidine), N2-(N,N-di-n-butyl-
formamidine), N2-phenylacetyl, N2-(1,2-diisobutyryloxyethylene), N2-(3-methoxy-4-
phenoxybenzoyl), N2-methoxyacetyl, chlorophenoxyacetyl, N2-isopropoxy-acetyl, N2-(2-
(tertbutyldiphenylsilyloxymethyl)-benzoyl), N2-phenylsulfonylethoxycarbonyl, N2-(4-
chlorophenylsulfonylethoxycarbonyl), N2-2-(acetoxymethyl) benzoyl, N.sup.2 -(3,4-
dichlorobenzoyl), O6 -Benzyl, O6-(2-(4-nitrophenyl)-ethyl), O6-(2-nitrophenyl), O6-(4-
nitrophenyl), O6-diphenylcarbamoyl, O6-(3,4-dimethoxybenzyl), O6-(3,5-dichlorophenyl),
O6-(2-cyanoethyl), O6-(2-trimethylsilylethyl), O6-phenylthioethyl, O6-(4-
nitrophenylthioethyl), O6-butylthiocarbonyl, O6-(6-methyl-3-pyridyl), O6-(2-(4-
nitrophenylsulfonyl)-ethyl), O6-(4-methyl-3-pyridyl), N2 -(4-nitrophenylethoxycarbonyl),
O6-allyl or any combination of these protecting groups for O6, N2-bis protection,


for Thymine:

O4-phenyl, O4-(2-(4-nitrophenyl)-ethyl), O4-(2-(4-nitro phenylsulfonyl)-ethyl), O4-(2-

methylphenyl), O4-(4-methyl phenyl), O4-(2,4-dimethylphenyl), N3 -benzoyl, N3 -(4-anisoyl), N3-(4-toluoyl), N3 -(2-toluoyl).

$R_1$ = 5'-protecting group such as:

9-Fluorenylmethoxycarbonyl, 4-chlorophenylsulfonylethoxy carbonyl, 4-nitrophenylsulfonyl-ethoxycarbonyl, phenyl sulfonylethoxycarbonyl, 2,2,2-tribromoethoxycarbonyl, levulinyl, 4,4',4"-tris(4,5-dichlorophtalimide)trityl, 4,4',4"-tris(benzoyloxy)trityl, 4,4',4"-tris(levulinyl oxy)trityl, p-anisyl-1-naphtylphenylmethyl, di-p-anisyl-1-naphtylmethyl, p-tolyldiphenylmethyl, 3-(imidazolylmethyl) -4,4'-dimethoxytrityl, methoxyacetyl, chloroacetyl, phenoxyacetyl, 4-chlorophenoxyacetyl, trityloxyacetyl, .beta.-benzoylpropionyl, isobutyloxycarbonyl, 4-nitrobenzyloxy carbonyl, 2-(methylthiomethoxymethyl)-benzoyl, 2-(iso propylthiomethoxymethyl) benzoyl, 4-(methylthiomethoxy butyryl, p-phenylazophenyloxycarbonyl, 2,4-dinitrophenyl ethoxycarbonyl, pivaloyl, 2-dibromomethylbenzoyl, tert-butyldimethylsilyl, 4,4'-dimethoxytrityl, 4'-monomethoxy trityl, 4-decyloxytrityl, 4-hexadecyloxytrityl, trityl, 1,1-bis-(4-methoxyphenyl)-1'-pyrenyl, 9-phenylxanthen-9-yl, 9-phenylthioxanthen-9-yl, 7-chloro-9-phenylthioxan then-9-yl, 9-(4-methoxyphenyl)-xanthen-9-yl, 9-(4-octadecyloxyphenyl)-xanthen-9-yl

$R_2$ = 3'-phosphate, 3'-(H-phosphonate), 3'-phosphoramidate, 3'-phosphoramidite, 3'-(alkanephosphonate) such as:

(a) 3'-phosphate: 2,2,2-Trichloroethyl, 2,2,2-tribromoethyl, 2-cyanoethyl, benzyl, 4-chlorophenyl, 4-nitrophenylethyl, 2-chlorophenyl, 2-diphenylmethylsilyl ethyl, phenylthio;

(b) 3'-phosphate esters: 2-Cyanoethyl-4-chlorophenyl, 2-cyanoethyl-2-cyanoethyl, 2-cyanoethyl-2-chlorophenyl, phenylsulfonylethyl-2-chlorophenyl, 9-fluorenylmethyl-2-chlorophenyl, 9-fluorenylmethyl-4-chlorophenyl, phenylsulfonylethyl-4-chlorophenyl, phenylsulfonylethyl-2-chloro phenyl, 2,2,2-tribromoethyl-4-chlorophenyl, 2,2,2-tribromo ethyl-2-chlorophenyl, 2,2,2-trichloroethyl-4-chlorophenyl, 2,2,2-trichloroethyl-2-chlorophenyl, 2-cyanoethyl-2-chloro -4-tritylphenyl, 2,2,2-tribromoethyl-2-chloro-4-tertbutyl phenyl, 4-nitrophenyl-phenyl, 2,4-dinitrobenzyl-2-chloro phenyl, 2,4-dinitrobenzyl-4-

chlorophenyl; S,S-diphenyl phosphorodithioate; 2-chlorophenyl-phosphoranilidate, phenyl-
phosphoranilidate,

(c) 3'-halophosphites (chloro or bromo): phenylsulfonylethoxy, methylsulfonylethoxy, 2-
(isopropyl sulfonyl)-ethoxy, 2-(tertbutylsulfonyl)-ethoxy, benzyl sulfonylethoxy, 4-
nitrobenzylsulfonylethoxy, 9-fluorenyl methoxy, 2-(4-nitrophenyl)-ethoxy, methoxy, 2-
cyano-1,1-dimethylethoxy, 2,2,2-trichloro-1,1-dimethylethoxy, 2,2,2-trichloroethoxy, 2-
cyanoethoxy, 2-cyano-1-methylethoxy, 2-cyano-1,1-dimethylethoxy-, 2-(4-nitrophenyl)-
ethoxy, 2(2-pyridyl)-ethoxy, 2-methylbenzyloxy, 4-chlorobenzyloxy, 2-chlorobenzyloxy,
2,4-dichlorobenzyloxy, 4-nitrobenzyloxy, allyloxy, phenoxy, 4-nitrophenoxy,
pentafluorophenyoxy, pentachlorophenoxy, 2,4,5-trichlorophenoxy, 2-bromophenoxy, 4-
bromophenoxy, 2-methylphenoxy, 2,6-dimethylphenoxy, 2,4-nitrophenoxy, 1,1,1,3,3,3-
hexafluoro-2-propyloxy, 2-chlorophenoxy.

(d) 3'-phosphoramidites: phenylsulfonylethoxydimethylamino, methylsulfonylethoxy-
morpholino, 2-(isopropylsulfonyl)-ethoxy-morpholino, 2-(tertbutylsulfonyl)-ethoxy-
morpholino, benzylsulfonylethoxy-morpholino, 4-nitrobenzylsulfonylethoxy-morpholino, 9-
fluorenylmethoxymorpholino, 2-(4-nitrophenyl)-ethoxy-morpholino, 2-(4-nitrophenyl)-
ethoxy-hexahydroazepine, 2-(4-nitrophenyl)ethoxy-octahydrazonine, 2-(4-nitrophenyl)-
ethoxyazacyclo tridecane, methoxy-pirrolidino, methoxy-piperidino, methoxy-diethylamino,
methoxy-diisopropilamino, methoxy-2,2,6,6-tetramethyl-N-piperidino, methoxy-morpholino,
2-cyano-1,1-dimethylethoxy-morpholino, 2,2,2-trichloro-1,1-dimethylethoxydimethylamino,
2,2,2-trichloro-1,1-dimethyl ethoxydiethylamino, 2,2,2-trichloro-1,1-dimethylethoxy-
diisopropylamino, 2,2,2-trichloro-1,1-dimethylethoxymorpholino, 2,2,2-
trichloroethoxydimethylamino, 2-cyanoethoxy diethylamino, 2-
cyanoethoxydiisopropylamino, 2-cyanoethoxy morpholino, 2-cyano-1-methylethoxy-
diethylamino, 2-cyano-1,1-dimethylethoxydiethylamino, 2-cyano-1,1-dimethylethoxy -
diisopropylamino, methylsulfonylethoxydiethylamino,
methylsulfonylethoxydiisopropylamino, 2,2,2-trichloroethoxydiisopropylamino, 2,2,2-
trichloro-1,1-dimethylethoxydiisopropylamino, 2-(4-nitrophenyl)-ethoxy-diisopropyl amino,
2(2-pyridyl)-ethoxy-diisopropylamino, 2(4-pyridyl)ethoxydiisopropylamino, 2-
methylbenzyloxy-diisopropyl amino, 4-chlorobenzyloxy-diisopropylamino, 2-chlorobenzyl
oxydiisopropylamino, 2,4-dichlorobenzyloxy-diisopropyl amino, 4-
nitrobenzyloxydiisopropylamino, allyloxydiiso propylamino, allyloxydimethylamino,
phenoxydiethylamino, 4-nitrophenoxydiethylamino, pentafluorophenoxydiethyl amino,

pentachlorophenoxy-diethylamino, 2,4,5-trichloro phenoxydiethylamino, 2-bromophenoxydiethylamino, 4-bromophenoxydiethylamino, 2-methylphenoxydiethylamino, 2,6-dimethylphenoxydiethylamino, 2,4-nitrophenoxy diethylamino, 1,1,1,3,3,3-hexafluoro-2-propyloxydiiso propylamino, 2-chlorophenoxy-morpholino, bis(diisopropyl amino), bis(diethylamino), bis(morpholino).

(e) phosphonate: methyl, ethyl, trifluoromethyl, cyanoethyl, trichloroethyl, tribromoethyl, trifluorethyl.

3. Therapeutic composition comprising the modified oligonucleotides and oligonucleosides according to claim 1 together with physiologically acceptable carriers.

4. A method for antisense therapy, comprising administration of the therapeutic composition according to claim 4 to a patient in need thereof.

5. A method according to claim 5, wherein the antisense therapy is against oncogenic sequences.

6. A method according to claim 5, wherein the antisense therapy is against pathogenic sequences.

7. A method according to claim 5, wherein the antisense therapy is for treatment of genetic disorders.

8. A diagnostic kit comprising the modified oligonucleotides and oligonucleosides according to claim 1.

9. A method of diagnosing nucleotide polymorphism(s) in an individual, comprising use of the diagnostic kit according to claim 8.

10. A DNA sequencing kit comprising the modified nucleosides and nucleotides.
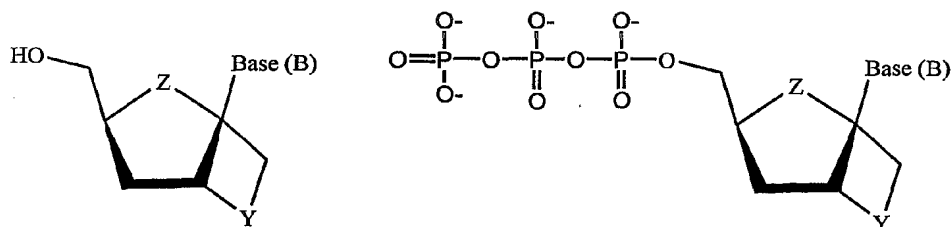


wherein combinations of modifications with Y, Z, or B are claimed:

Y = O, S, or NH or $NCH_3$, $CH_2$ or $CH(CH_3)$;
Z = O, S, or NH or $NCH_3$, $CH_2$ or $CH(CH_3)$
B = A, C, G, T, U, 5-F/Cl/Br-U; 7-deaza-G or hypoxanthine

α– or β–D-(or L) ribo, xylo , arabino or lyxo configuration

11. Use of the nucleotides and nucleosides according to claim 1 for production of aptamers.

12. Use of the compounds according to claims 1, 2 and/or 10 for drug development .

13. Use of the compounds according to claims 1, 2 and/or 10 in any form of polymerase chain reaction (PCR).

14. Use of the compounds according to claims 1, 2 and/or 10 in any molecular biology kit for diagnosis, detection or as reagent.

1/2

FIG 1
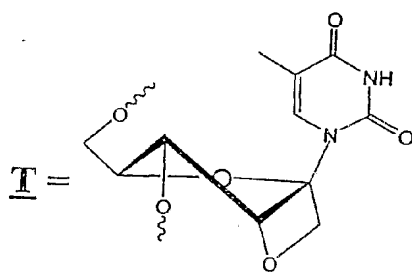
# FIG. 2
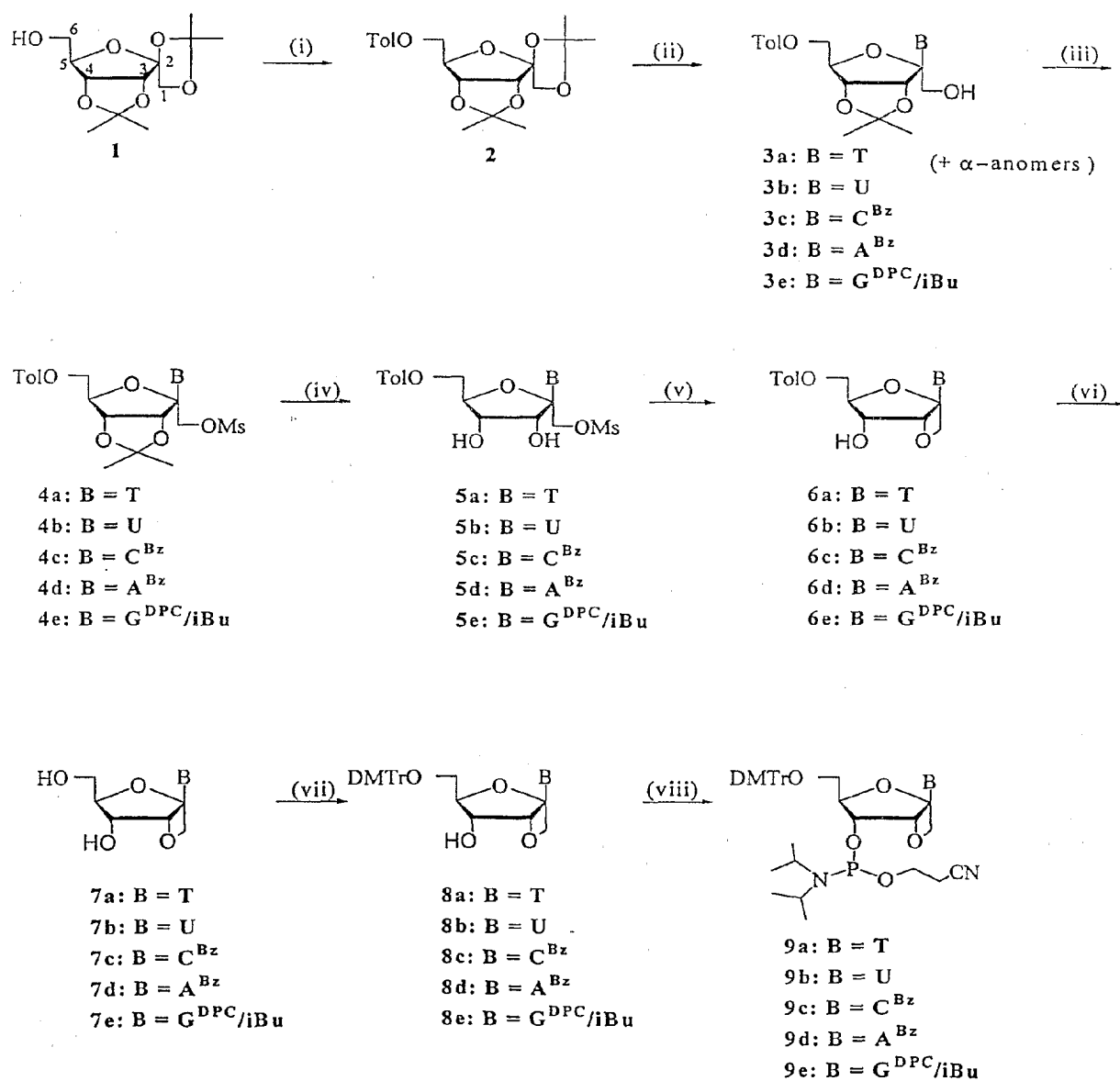


3a: B = T       (+ α−anomers)
3b: B = U
3c: B = C$^{Bz}$
3d: B = A$^{Bz}$
3e: B = G$^{DPC}$/iBu

4a: B = T
4b: B = U
4c: B = C$^{Bz}$
4d: B = A$^{Bz}$
4e: B = G$^{DPC}$/iBu

5a: B = T
5b: B = U
5c: B = C$^{Bz}$
5d: B = A$^{Bz}$
5e: B = G$^{DPC}$/iBu

6a: B = T
6b: B = U
6c: B = C$^{Bz}$
6d: B = A$^{Bz}$
6e: B = G$^{DPC}$/iBu

7a: B = T
7b: B = U
7c: B = C$^{Bz}$
7d: B = A$^{Bz}$
7e: B = G$^{DPC}$/iBu

8a: B = T
8b: B = U
8c: B = C$^{Bz}$
8d: B = A$^{Bz}$
8e: B = G$^{DPC}$/iBu

9a: B = T
9b: B = U
9c: B = C$^{Bz}$
9d: B = A$^{Bz}$
9e: B = G$^{DPC}$/iBu

**SUBSTITUTE SHEET (RULE 26)**

# INTERNATIONAL SEARCH REPORT

## A. CLASSIFICATION OF SUBJECT MATTER

IPC7: C07H 7/06, A61K 31/7052, C12P 19/30, C07H 19/00, C07F 9/02

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC7: C07H, C12P, C07F, A61K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

SE,DK,FI,NO classes as above

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

| Category* | Citation of document, with indication, where appropriate, of the relevant passages | Relevant to claim No. |
|---|---|---|
| P,X | J. Chem. Soc., Perkin Trans., Vol. 2, 2001, Pushpangadan I. Pradeepkumar et al: "Conformation-Specific cleavage of antisense oligonucleotide-RNA duplexes by RNase H", page 402 - page 408, scheme 1 | 1,10, PART OF CLAIMS 2,12-14 |
| | -- | |
| X | Tetrahedron Letters, Vol. 41, September 2000, P. I. Pradeepkumar etal: "Transmission of the conformational information in the antisense/RNA hybrid duplex influences the patterna of the RNase H cleavagne reaction", page 8601 - page 8607 | 1,10, PART OF CLAIMS 2,12-14 |
| | -- | |

[X] Further documents are listed in the continuation of Box C.    [X] See patent family annex.

| * | Special categories of cited documents: |
|---|---|
| "A" | document defining the general state of the art which is not considered to be of particular relevance |
| "E" | earlier application or patent but published on or after the international filing date |
| "L" | document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified) |
| "O" | document referring to an oral disclosure, use, exhibition or other means |
| "P" | document published prior to the international filing date but later than the priority date claimed |

| "T" | later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention |
|---|---|
| "X" | document of particular relevance: the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone |
| "Y" | document of particular relevance: the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art |
| "&" | document member of the same patent family |

| Date of the actual completion of the international search | Date of mailing of the international search report |
|---|---|
| 28 March 2002 | 0 3 -04- 2002 |

| Name and mailing address of the ISA/ | Authorized officer |
|---|---|
| Swedish Patent Office Box 5055, S-102 42 STOCKHOLM | Fernando Farieta/BS |
| Facsimile No. + 46 8 666 02 86 | Telephone No. + 46 8 782 25 00 |

Form PCT/ISA/210 (second sheet) (July 1998)

C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT

| Category* | Citation of document, with indication, where appropriate, of the relevant passages | Relevant to claim No. |
|---|---|---|
| X | Collection Czechosiov. Chem. Commun., Vol. 39, 1997, H. Hrebabeckýet al: "Synthesis of 1-beta-D-Psicofuranosyluracil and 1-beta-D-psicofuranosylcytosine", page 1098 - page 1106, figure XIIIa, XVa, XVIa | 1,10, PART OF CLAIMS 2,12-14 |
| X | Nucleosides & Nucleotides, Volume 11, No. 1, 1992, Robert D. Elliott et al: "Synthesis of 9-(1-Deoxy-1-Phosphone-Beta-D-Psicofuranosyl)-1,9-(Dihydro-6H-Purin-6-One As Potential Transition State Analog Inhibitor of Purine Nucleoside Phosphorylase", page 97-119, page 99 | 1,10, PART OF CLAIMS 2,12-14 |
| A | US 3126372 A (BANNISTER), 24 March 1964 (24.03.64) | 1,10, PART OF CLAIMS 2,12-14 |
| A | US 3125567 A (WILLIAM SCHROEDER), 17 March 1964 (17.03.64) | 1,10, PART OF CLAIMS 2,12-14 |
| A | US 3079378 A (WILLIAM SCHROEDER), 26 February 1963 (26.02.63) | 1,10, PART OF CLAIMS 2,12-14 |
| A | US 3020274 A (THOMAS E. EBLE ET AL), 6 February 1962 (06.02.62) | 1,10, PART OF CLAIMS 2,12-14 |

# INTERNATIONAL SEARCH REPORT

International application No.
PCT/SE01/02484

| Box I | Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet) |
|---|---|

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:

2. ☒ Claims Nos.: **3-9**
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:

**see next sheet***

3. ☐ Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

| Box II | Observations where unity of invention is lacking (Continuation of item 2 of first sheet) |
|---|---|

This International Searching Authority found multiple inventions in this international application, as follows:

**see next sheet****

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.

2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.

3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:

4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

**Remark on Protest**   ☐ The additional search fees were accompanied by the applicant's protest.

☐ No protest accompanied the payment of additional search fees.

Form PCT/ISA/210 (continuation of first sheet (1)) (July1998)

\*

Claims 3-9 relate to methods of treatment of the human or animal body by surgery or by therapy/ diagnostic methods practised on the human or animal body/ Rule. 39.1.(iv). Nevertheless, a search has been executed for these claims. (Claims 4 and 5 present obvious errors according to PCT Rule 91)

\*\*

According to PCT Rule 13.2, an international application shall relate to one invention only or a group of inventions linked by one or more of the same or corresponding special technical features, i. e. features that define a contribution which each of the inventions makes over the prior art. The present application is thus considered to relate to at least five inventions, namely:

1) Invention 1: Modified nucleosides and nucleotides and use thereof according to claims 1, 10 and part of claims 12-14.

2) Invention 2: Reagents for the preparation of modified nucleosides-nucleotides analogs, namely reagent for thymine, according to claim 2 and part of claims 12-14.

3) Invention 3: Reagents for the preparation of modified nucleosides-nucleotides analogs, namely reagent for adenine, according to claim 2 and part of claims 12-14

4) Invention 4: Reagents for the preparation of modified nucleosides-nucleotides analogs, namely reagent for cytosine, according to claim 2 and part of claims 12-14

5) Invention 5: Reagents for the preparation of modified nucleosides-nucleotides analogs, namely reagent for uracil, according to claim 2 and part of claims 12-14

The use of a reagent has been shown only for the case of the reagent "toluoyl" in the description; it gives support partially to invention 2. Consequently, the use of the reagents claimed in claim 2 has not been performed for most of the reagent variants; this prerequisite is not fulfilled in the present application.

Inventions 1 and 2 have been partially searched.

| Patent document cited in search report | | | Publication date | Patent family member(s) | Publication date |
|---|---|---|---|---|---|
| US | 3126372 | A | 24/03/64 | NONE | |
| US | 3125567 | A | 17/03/64 | NONE | |
| US | 3079378 | A | 26/02/63 | NONE | |
| US | 3020274 | A | 06/02/62 | NONE | |